## SEQUENCE LISTING

<110>	The University of North Carolina Threadgill, David W Lee, Daekee	
<120>	MODULATION OF EPIDERMAL GROWTH FACTOR HETERODIMER ACTIVITY	
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Leu Glu Ile Thr Tyr Val Gln Arg Asn Tyr Asp Leu Ser Phe Leu Lys 70 75 80

Thr Ile Gln Glu Val Ala Gly Tyr Val Leu Ile Ala Leu Asn Thr Val 85 90 95

Glu Arg Ile Pro Leu Glu Asn Leu Gln Ile Ile Arg Gly Asn Met Tyr 100 105 110

Tyr Glu Asn Ser Tyr Ala Leu Ala Val Leu Ser Asn Tyr Asp Ala Asn 115 120 125

Lys Thr Gly Leu Lys Glu Leu Pro Met Arg Asn Leu Gln Glu Ile Leu 130 135 140

His Gly Ala Val Arg Phe Ser Asn Asn Pro Ala Leu Cys Asn Val Glu 145 150 155 160

Ser Ile Gln Trp Arg Asp Ile Val Ser Ser Asp Phe Leu Ser Asn Met 165 170 175

Ser Met Asp Phe Gln Asn His Leu Gly Ser Cys Gln Lys Cys Asp Pro 180 185 190

Ser Cys Pro Asn Gly Ser Cys Trp Gly Ala Gly Glu Asn Cys Gln 195 200 205

Lys Leu Thr Lys Ile Ile Cys Ala Gln Gln Cys Ser Gly Arg Cys Arg 210 215 220

Gly Lys Ser Pro Ser Asp Cys Cys His Asn Gln Cys Ala Ala Gly Cys 225 230 235 240

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Glu Ala Thr Cys Lys Asp Thr Cys Pro Pro Leu Met Leu Tyr Asn Pro 260 265 270

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Gly Ser Cys Val Arg Ala Cys Gly Ala Asp Ser Tyr Glu Met Glu Glu 305 310 315 320

Asp Gly Val Arg Lys Cys Lys Cys Glu Gly Pro Cys Arg Lys Val 325 330 335

Cys Asn Gly Ile Gly Ile Gly Glu Phe Lys Asp Ser Leu Ser Ile Asn  $340 \hspace{1.5cm} 345 \hspace{1.5cm} 350$ 

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Leu His Ile Leu Pro Val Ala Phe Arg Gly Asp Ser Phe Thr His Thr 370 375 380

Pro Pro Leu Asp Pro Gln Glu Leu Asp Ile Leu Lys Thr Val Lys Glu 385 390 395 400

Ile Thr Gly Phe Leu Leu Ile Gln Ala Trp Pro Glu Asn Arg Thr Asp 405 410 415

Leu His Ala Phe Glu Asn Leu Glu Ile Ile Arg Gly Arg Thr Lys Gln 420 425 430

His Gly Gln Phe Ser Leu Ala Val Val Ser Leu Asn Ile Thr Ser Leu 435 440 445

Gly Leu Arg Ser Leu Lys Glu Ile Ser Asp Gly Asp Val Ile Ile Ser 450 455 460

Gly Asn Lys Asn Leu Cys Tyr Ala Asn Thr Ile Asn Trp Lys Lys Leu 465 470 480

Phe Gly Thr Ser Gly Gln Lys Thr Lys Ile Ile Ser Asn Arg Gly Glu 485 490 495

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- Lys Val Lys Ile Pro Val Ala Ile Lys Glu Leu Arg Glu Ala Thr Ser 740 Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Ser Val Asp Asn Pro His Val Cys Arg Leu Gly Ile Cys Leu Thr Ser Thr Val Gln Leu Ile Thr Gln Leu Met Pro Phe Gly Cys Leu Leu Asp 790 795 Tyr Val Arg Glu His Lys Asp Asn Ile Gly Ser Gln Tyr Leu Leu Asn Trp Cys Val Gln Ile Ala Lys Gly Met Asn Tyr Leu Glu Asp Arg Arg 825 Leu Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys Thr Pro 840 Gln His Val Lys Ile Thr Asp Phe Gly Leu Ala Lys Leu Leu Gly Ala 855 Glu Glu Lys Glu Tyr His Ala Glu Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu Glu Ser Ile Leu His Arg Ile Tyr Thr His Gln Ser Asp 890 895 Val Trp Ser Tyr Gly Val Thr Val Trp Glu Leu Met Thr Phe Gly Ser 900 905 910
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920

915

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- Asp Thr Phe Leu Pro Val Pro Glu Tyr Ile Asn Gln Ser Val Pro 1085 1090 1095
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- Pro Leu Asn Pro Ala Pro Ser Arg Asp Pro His Tyr Gln Asp Pro 1115 1120 1125
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- Pro Thr Cys Val Asn Ser Thr Phe Asp Ser Pro Ala His Trp Ala 1145 1150 1155
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<sup>&</sup>lt;213> Mus musculus

<sup>&</sup>lt;400> 4

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Glu Ala Thr Cys Lys Asp Thr Cys Pro Pro Leu Met Leu Tyr Asn Pro Thr Thr Tyr Gln Met Asp Val Asn Pro Glu Gly Lys Tyr Ser Phe Gly Ala Thr Cys Val Lys Lys Cys Pro Arg Asn Tyr Val Val Thr Asp His Gly Ser Cys Val Arg Ala Cys Gly Pro Asp Tyr Tyr Glu Val Glu Glu Asp Gly Ile Arg Lys Cys Lys Cys Asp Gly Pro Cys Arg Lys Val Cys Asn Gly Ile Gly Ile Gly Glu Phe Lys Asp Thr Leu Ser Ile Asn Ala Thr Asn Ile Lys His Phe Lys Tyr Cys Thr Ala Ile Ser Gly Asp Leu His Ile Leu Pro Val Ala Phe Lys Gly Asp Ser Phe Thr Arg Thr Pro Pro Leu Asp Pro Arg Glu Leu Glu Ile Leu Lys Thr Val Lys Glu Ile Thr Gly Phe Leu Leu Ile Gln Ala Trp Pro Asp Asn Trp Thr Asp Leu His Ala Phe Glu Asn Leu Glu Ile Ile Arg Gly Arg Thr Lys Gln His Gly Gln Phe Ser Leu Ala Val Val Gly Leu Asn Ile Thr Ser Leu Gly Leu Arg Ser Leu Lys Glu Ile Ser Asp Gly Asp Val Ile Ile Ser Gly Asn Arg Asn Leu Cys Tyr Ala Asn Thr Ile Asn Trp Lys Lys Leu Phe Gly Thr Pro Asn Gln Lys Thr Lys Ile Met Asn Asn Arg Ala Glu

Lys Asp Cys Lys Ala Val Asn His Val Cys Asn Pro Leu Cys Ser Ser Glu Gly Cys Trp Gly Pro Glu Pro Arg Asp Cys Val Ser Cys Gln Asn Val Ser Arg Gly Arg Glu Cys Val Glu Lys Cys Asn Ile Leu Glu Gly Glu Pro Arg Glu Phe Val Glu Asn Ser Glu Cys Ile Gln Cys His Pro Glu Cys Leu Pro Gln Ala Met Asn Ile Thr Cys Thr Gly Arg Gly Pro Asp Asn Cys Ile Gln Cys Ala His Tyr Ile Asp Gly Pro His Cys Val Lys Thr Cys Pro Ala Gly Ile Met Gly Glu Asn Asn Thr Leu Val Trp Lys Tyr Ala Asp Ala Asn Asn Val Cys His Leu Cys His Ala Asn Cys Thr Tyr Gly Cys Ala Gly Pro Gly Leu Gln Gly Cys Glu Val Trp Pro Ser Gly Pro Lys Ile Pro Ser Ile Ala Thr Gly Ile Val Gly Leu Leu Phe Ile Val Val Val Ala Leu Gly Ile Gly Leu Phe Met Arg Arg Arg His Ile Val Arg Lys Arg Thr Leu Arg Arg Leu Leu Gln Glu Arg Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Glu Ala Pro Asn Gln Ala His Leu Arg Ile Leu Lys Glu Thr Glu Phe Lys Lys Ile Lys Val Leu Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys Gly Leu Trp Ile Pro Glu

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7	ľhr	Ser	Pro 755	Lys	Ala	Asn	Lys	Glu 760	Ile	Leu	Asp	Glu	Ala 765	Tyr	Val	Met
I	Ala	Ser 770	Val	Asp	Asn	Pro	His 775	Val	Cys	Arg	Leu	Leu 780	Gly	Ile	Cys	Leu
	Thr 785	Ser	Thr	Val	Gln	Leu 790	Ile	Thr	Gln	Leu	Met 795	Pro	Tyr	Gly	Cys	Leu 800
Ι	Leu	Asp	Tyr	Val	Arg 805	Glu	His	Lys	Asp	Asn 810	Ile	Gly	Ser	Gln	Tyr 815	Leu
Ι	Leu	Asn	Trp	Cys 820	Val	Gln	Ile	Ala	Lys 825	Gly	Met	Asn	Tyr	Leu 830	Glu	Asp
I	Arg	Arg	Leu 835	Val	His	Arg	Asp	Leu 840	Ala	Ala	Arg	Asn	Val 845	Leu	Val	Lys
1	hr	Pro 850	Gln	His	Val	Lys	Ile 855	Thr	Asp	Phe	Gly	Leu 860	Ala	Lys	Leu	Leu

Lys Trp Met Ala Leu Glu Ser Ile Leu His Arg Ile Tyr Thr His Gln 885 890 895

Gly Ala Glu Glu Lys Glu Tyr His Ala Glu Gly Gly Lys Val Pro Ile

875

870

865

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Gly Ser Lys Pro Tyr Asp Gly Ile Pro Ala Ser Asp Ile Ser Ser Ile 915 920 925

Leu Glu Lys Gly Glu Arg Leu Pro Gln Pro Pro Ile Cys Thr Ile Asp 930 935 940

Val Tyr Met Ile Met Val Lys Cys Trp Met Ile Asp Ala Asp Ser Arg 945 950 955 960

- Pro Lys Phe Arg Glu Leu Ile Leu Glu Phe Ser Lys Met Ala Arg Asp 965 970 975
- Pro Gln Arg Tyr Leu Val Ile Gln Gly Asp Glu Arg Met His Leu Pro 980 985 990
- Ser Pro Thr Asp Ser Asn Phe Tyr Arg Ala Leu Met Asp Glu Glu Asp 995 1000 1005
- Met Glu Asp Val Val Asp Ala Asp Glu Tyr Leu Ile Pro Gln Gln 1010 1015 1020
- Gly Phe Phe Asn Ser Pro Ser Thr Ser Arg Thr Pro Leu Leu Ser 1025 1030 1035
- Ser Leu Ser Ala Thr Ser Asn Asn Ser Thr Val Ala Cys Ile Asn 1040 1045 1050
- Arg Asn Gly Ser Cys Arg Val Lys Glu Asp Ala Phe Leu Gln Arg 1055 1060 1065
- Tyr Ser Ser Asp Pro Thr Gly Ala Val Thr Glu Asp Asn Ile Asp 1070 1075 1080
- Asp Ala Phe Leu Pro Val Pro Glu Tyr Val Asn Gln Ser Val Pro 1085 1090 1095
- Lys Arg Pro Ala Gly Ser Val  $\mbox{Gln}$  Asn Pro Val  $\mbox{Tyr}$   $\mbox{His}$  Asn  $\mbox{Gln}$   $\mbox{1100}$
- Pro Leu His Pro Ala Pro Gly Arg Asp Leu His Tyr Gln Asn Pro 1115 1120 1125
- His Ser Asn Ala Val Gly Asn Pro Glu Tyr Leu Asn Thr Ala Gln 1130 1135 1140
- Pro Thr Cys Leu Ser Ser Gly Phe Asn Ser Pro Ala Leu Trp Ile 1145 1150 1155
- Gln Lys Gly Ser His Gln Met Ser Leu Asp Asn Pro Asp Tyr Gln 1160 1165 1170
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Gly Pro Thr Ala Glu Asn Ala Glu Tyr Leu Arg Val Ala Pro Pro

1200

1195

1190

115

110

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			gtc Val									823
			tgc Cys					-	_			871
	-		 gga Gly	 		_	_		_			919
			tgt Cys									967
			caa Gln 235									1015
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					Gly ggg												1687
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gag Glu	tgc Cys 535	gtg Val	gag Glu	aaa Lys	tgc Cys	aac Asn 540	atc Ile	ctg Leu	gag Glu	Gly	gaa Glu 545	cca Pro	agg Arg	gag Glu	ttt Phe		1927
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					atg Met 795											2	695
					att Ile											2	2743
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													gtc Val	2839
													aaa Lys	2887
													ttg Leu 885	2935
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													gag Glu 965	3175
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- Leu Ser Leu Gln Arg Met Tyr Asn Asn Cys Glu Val Val Leu Gly Asn 50 55 60
- Leu Glu Ile Thr Tyr Val Gln Arg Asn Tyr Asp Leu Ser Phe Leu Lys 70 75 80
- Thr Ile Gln Glu Val Ala Gly Tyr Val Leu Ile Ala Leu Asn Thr Val 85 90 95
- Glu Arg Ile Pro Leu Glu Asn Leu Gln Ile Ile Arg Gly Asn Ala Leu 100 105 110
- Tyr Glu Asn Thr Tyr Ala Leu Ala Ile Leu Ser Asn Tyr Gly Thr Asn 115 120 125
- Arg Thr Gly Leu Arg Glu Leu Pro Met Arg Asn Leu Gln Glu Ile Leu 130 135 140
- Ile Gly Ala Val Arg Phe Ser Asn Asn Pro Ile Leu Cys Asn Met Asp 145 150 155 160
- Thr Ile Gln Trp Arg Asp Ile Val Gln Asn Val Phe Met Ser Asn Met 165 170 175
- Ser Met Asp Leu Gln Ser His Pro Ser Ser Cys Pro Lys Cys Asp Pro 180 185 190
- Ser Cys Pro Asn Gly Ser Cys Trp Gly Gly Gly Glu Glu Asn Cys Gln 195 200 205
- Lys Leu Thr Lys Ile Ile Cys Ala Gln Gln Cys Ser His Arg Cys Arg 210 215 220
- Gly Arg Ser Pro Ser Asp Cys Cys His Asn Gln Cys Ala Ala Gly Cys 225 230 235 240
- Thr Gly Pro Arg Glu Ser Asp Cys Leu Val Cys Gln Lys Phe Gln Asp 245 250 255

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Thr Thr Tyr Gln Met Asp Val Asn Pro Glu Gly Lys Tyr Ser Phe Gly 275 280 285

Ala Thr Cys Val Lys Lys Cys Pro Arg Asn Tyr Val Val Thr Asp His 290 295 300

Gly Ser Cys Val Arg Ala Cys Gly Pro Asp Tyr Tyr Glu Val Glu Glu 305 310 315 320

Asp Gly Ile Arg Lys Cys Lys Cys Asp Gly Pro Cys Arg Lys Val 325 330 335

Cys Asn Gly Ile Gly Ile Gly Glu Phe Lys Asp Thr Leu Ser Ile Asn 340 345 350

Ala Thr Asn Ile Lys His Phe Lys Tyr Cys Thr Ala Ile Ser Gly Asp 355 360 365

Leu His Ile Leu Pro Val Ala Phe Lys Gly Asp Ser Phe Thr Arg Thr 370 375 380

Pro Pro Leu Asp Pro Arg Glu Leu Glu Ile Leu Lys Thr Val Lys Glu 385 390 395 400

Ile Thr Gly Phe Leu Leu Ile Gln Ala Trp Pro Asp Asn Trp Thr Asp 405 410 415

Leu His Ala Phe Glu Asn Leu Glu Ile Ile Arg Gly Arg Thr Lys Gln
420 425 430

His Gly Gln Phe Ser Leu Ala Val Val Gly Leu Asn Ile Thr Ser Leu 435 440 445

Gly Leu Arg Ser Leu Lys Glu Ile Ser Asp Gly Asp Val Ile Ile Ser 450 455 460

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Phe Gly Thr Pro Asn Gln Lys Thr Lys Ile Met Asn Asn Arg Ala Glu 485 490 495 Lys Asp Cys Lys Ala Val Asn His Val Cys Asn Pro Leu Cys Ser Ser 500 505 510

Glu Gly Cys Trp Gly Pro Glu Pro Arg Asp Cys Val Ser Cys Gln Asn 515 520 525

Val Ser Arg Gly Arg Glu Cys Val Glu Lys Cys Asn Ile Leu Glu Gly 530 540

Glu Pro Arg Glu Phe Val Glu Asn Ser Glu Cys Ile Gln Cys His Pro 545 550 555 560

Glu Cys Leu Pro Gln Ala Met Asn Ile Thr Cys Thr Gly Arg Gly Pro 565 570 575

Asp Asn Cys Ile Gln Cys Ala His Tyr Ile Asp Gly Pro His Cys Val580  $\phantom{0}585$   $\phantom{0}590$ 

Lys Thr Cys Pro Ala Gly Ile Met Gly Glu Asn Asn Thr Leu Val Trp  $595 \hspace{1.5cm} 600 \hspace{1.5cm} 605$ 

Lys Tyr Ala Asp Ala Asn Asn Val Cys His Leu Cys His Ala Asn Cys 610 620

Thr Tyr Gly Cys Ala Gly Pro Gly Leu Gln Gly Cys Glu Val Trp Pro 625 630 635 640

Ser Gly Pro Lys Ile Pro Ser Ile Ala Thr Gly Ile Val Gly Leu 645 650 655

Leu Phe Ile Val Val Val Ala Leu Gly Ile Gly Leu Phe Met Arg Arg 660 665 670

Arg His Ile Val Arg Lys Arg Thr Leu Arg Arg Leu Leu Gln Glu Arg 675 680 685

Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Glu Ala Pro Asn Gln Ala 690 695 700

His Leu Arg Ile Leu Lys Glu Thr Glu Phe Lys Lys Ile Lys Val Leu 705 710 715 720

Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys Gly Leu Trp Ile Pro Glu
725 730 735

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740 745 750

Thr Ser Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met 755 760 765

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Thr Ser Thr Val Gln Leu Ile Thr Gln Leu Met Pro Tyr Gly Cys Leu 785 790 795 800

Leu Asp Tyr Val Arg Glu His Lys Asp Asn Ile Gly Ser Gln Tyr Leu 805 810 815

Leu Asn Trp Cys Val Gln Ile Ala Lys Gly Met Asn Tyr Leu Glu Asp 820 825 830

Arg Arg Leu Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys 835 840 845

Thr Pro Gln His Val Lys Ile Thr Gly Phe Gly Leu Ala Lys Leu Leu 850 855 860

Gly Ala Glu Glu Lys Glu Tyr His Ala Glu Gly Gly Lys Val Pro Ile 865 870 875 880

Lys Trp Met Ala Leu Glu Ser Ile Leu His Arg Ile Tyr Thr His Gln 885 890 895

Ser Asp Val Trp Ser Tyr Gly Val Thr Val Trp Glu Leu Met Thr Phe 900 905 910

Gly Ser Lys Pro Tyr Asp Gly Ile Pro Ala Ser Asp Ile Ser Ser Ile 915 920 925

Leu Glu Lys Gly Glu Arg Leu Pro Gln Pro Pro Ile Cys Thr Ile Asp 930 935 940

Val Tyr Met Ile Met Val Lys Cys Trp Met Ile Asp Ala Asp Ser Arg 945 950 955 960

Pro Lys Phe Arg Glu Leu Ile Leu Glu Phe Ser Lys Met Ala Arg Asp 965 970 975

- Pro Gln Arg Tyr Leu Val Ile Gln Gly Asp Glu Arg Met His Leu Pro 980 985 990
- Ser Pro Thr Asp Ser Asn Phe Tyr Arg Ala Leu Met Asp Glu Glu Asp 995 1000 1005
- Met Glu Asp Val Val Asp Ala Asp Glu Tyr Leu Ile Pro Gln Gln 1010 1015 1020
- Gly Phe Phe Asn Ser Pro Ser Thr Ser Arg Thr Pro Leu Leu Ser 1025 1030 1035
- Ser Leu Ser Ala Thr Ser Asn Asn Ser Thr Val Ala Cys Ile Asn 1040 1045 1050
- Arg Asn Gly Ser Cys Arg Val Lys Glu Asp Ala Phe Leu Gln Arg 1055 1060 1065
- Tyr Ser Ser Asp Pro Thr Gly Ala Val Thr Glu Asp Asn Ile Asp 1070 1075 1080
- Asp Ala Phe Leu Pro Val Pro Glu Tyr Val Asn Gln Ser Val Pro 1085 1090 1095
- Lys Arg Pro Ala Gly Ser Val Gln Asn Pro Val Tyr His Asn Gln 1100 1105 1110
- Pro Leu His Pro Ala Pro Gly Arg Asp Leu His Tyr Gln Asn Pro 1115 1120 1125
- His Ser Asn Ala Val Gly Asn Pro Glu Tyr Leu Asn Thr Ala Gln 1130 1135 1140
- Pro Thr Cys Leu Ser Ser Gly Phe Asn Ser Pro Ala Leu Trp Ile 1145 1150 1155
- Gln Lys Gly Ser His Gln Met Ser Leu Asp Asn Pro Asp Tyr Gln 1160 1165 1170
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cccacgo	gggc cetttactge geegegegee eggeeeeeae eeetegeage aeeeegegee	180
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	g ctg gcg gcc ttg tgc cgc tgg ggg ctc ctc	286
ccc ccc Pro Pro	c gga gcc gcg agc acc caa gtg tgc acc ggc aca gac atg aag o Gly Ala Ala Ser Thr Gln Val Cys Thr Gly Thr Asp Met Lys 20 25 30	334

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		_		_	cag Gln			-			_	_				430
					agc Ser 70											478
					atc Ile											526
					gtg Val											574
					gac Asp											622
					cca Pro											670
ctc Leu 145	aca Thr	gag Glu	atc Ile	ttg Leu	aaa Lys 150	gga Gly	Gl <sup>A</sup> aaa	gtc Val	ttg Leu	atc Ile 155	cag Gln	cgg Arg	aac Asn	ccc Pro	cag Gln 160	718
					acg Thr											766
					aca Thr											814
					atg Met											862
tct Ser	gag Glu 210	gat Asp	tgt Cys	cag Gln	agc Ser	ctg Leu 215	acg Thr	cgc Arg	act Thr	gtc Val	tgt Cys 220	gcc Ala	ggt Gly	ggc Gly	tgt Cys	910
gcc Ala 225	cgc Arg	tgc Cys	aag Lys	ggg Gly	cca Pro 230	ctg Leu	ccc Pro	act Thr	gac Asp	tgc Cys 235	tgc Cys	cat His	gag Glu	cag Gln	tgt Cys 240	958
					ggc Gly											1006
cac His	ttc Phe	aac Asn	cac His 260	agt Ser	ggc Gly	atc Ile	tgt Cys	gag Glu 265	ctg Leu	cac His	tgc Cys	cca Pro	gcc Ala 270	ctg Leu	gtc Val	1054

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					agt Ser											1342
aag Lys	atc Ile 370	ttt Phe	GJ À GG À	agc Ser	ctg Leu	gca Ala 375	ttt Phe	ctg Leu	ccg Pro	gag Glu	agc Ser 380	ttt Phe	gat Asp	ggg	gac Asp	1390
					gcc Ala 390											1438
					atc Ile											1486
					ctc Leu											1534
					aat Asn											1582
ggc Gly	atc Ile 450	agc Ser	tgg Trp	ctg Leu	Gly	ctg Leu 455	cgc Arg	tca Ser	ctg Leu	agg Arg	gaa Glu 460	ctg Leu	ggc Gly	agt Ser	gga Gly	1630
					cat His 470											1678
					ttt Phe											1726
					gac Asp											1774

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					ttc Phe											18	370
					ctc Leu 550											19	918
					gag Glu											19	966
					gac Asp											20	14
					gcc Ala											20	62
					tgg Trp											21	.10
					tgc Cys 630											21	.58
					cag Gln											22	206
					ctg Leu											22	254
					cgg Arg											23	302
					acg Thr											23	50
					gcg Ala 710											23	98
					ctt Leu											24	46
ggc Gly	atc Ile	tgg Trp	atc Ile 740	cct Pro	gat Asp	ggg Gly	gag Glu	aat Asn 745	gtg Val	aaa Lys	att Ile	cca Pro	gtg Val 750	gcc Ala	atc Ile	24	94

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	tac Tyr								2590
	atc Ile								2638
	ggc Gly								2686
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	ctg Leu								2782
	ctg Leu	_	_		_			_	2830
	cgg Arg								2878
	gtg Val								2926
	acc Thr 900								2974
	atg Met								3022
	cct Pro								3070
	acc Thr								3118
	gaa Glu								3166
	gcc Ala 980								3214

_	Leu		-	_	Pro I	-	-	_		ttc t Phe T 1		-		_	3262
		Asp				Āsp				gct Ala 1020	Glu				3307
		Pro				Phe				cct Pro 1035	Ala				3352
						Arg				tca Ser 1050	Ser	acc Thr			3397
						Leu				ccc Pro 1065	Ser	gaa Glu			3442
						Ala				ggg Gly 1080	Ala				3487
						Gly				gcc Ala 1095	Lys				3532
	agc Ser 1100					Pro				cag Gln 1110	Arg	tac Tyr	_		3577
	gac Asp 1115					Pro				gat Asp 1125	Gly				3622
	ccc Pro 1130					Gln				gtg Val 1140	Asn	cag Gln			3667
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						Thr				ccc Pro 1170					3757
						Val				ttt Phe 1185	Ala				3802
	gcc Ala 1190					Tyr				cag Gln 1200					3847
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Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His 35 40 45

Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr 50 55 60

Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val 65 70 75 80

Gln Gly Tyr Val Leu Ile Ala His Asn Gln Val Arg Gln Val Pro Leu 85 90 95

Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Asn Tyr 100 105 110

Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Asn Asn Thr Thr Pro 115 120 125

Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln Leu Arg Ser 130 135 140

Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Gln Arg Asn Pro Gln 145 150 155 160

Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe His Lys Asn 165 170 175

Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser Arg Ala Cys 180 185 190

His Pro Cys Ser Pro Met Cys Lys Gly Ser Arg Cys Trp Gly Glu Ser 195 200 205

Ser Glu Asp Cys Gln Ser Leu Thr Arg Thr Val Cys Ala Gly Gly Cys 210 215 220

Ala Arg Cys Lys Gly Pro Leu Pro Thr Asp Cys Cys His Glu Gln Cys 225 230 235 240

Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu Ala Cys Leu 245 250 255

His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro Ala Leu Val 260 265 270

Thr Tyr Asn Thr Asp Thr Phe Glu Ser Met Pro Asn Pro Glu Gly Arg 275 280 285

Tyr Thr Phe Gly Ala Ser Cys Val Thr Ala Cys Pro Tyr Asn Tyr Leu 290 295 300

Ser Thr Asp Val Gly Ser Cys Thr Leu Val Cys Pro Leu His Asn Gln 305 310 315 320

Glu Val Thr Ala Glu Asp Gly Thr Gln Arg Cys Glu Lys Cys Ser Lys Pro Cys Ala Arg Val Cys Tyr Gly Leu Gly Met Glu His Leu Arg Glu Val Arg Ala Val Thr Ser Ala Asn Ile Gln Glu Phe Ala Gly Cys Lys Lys Ile Phe Gly Ser Leu Ala Phe Leu Pro Glu Ser Phe Asp Gly Asp Pro Ala Ser Asn Thr Ala Pro Leu Gln Pro Glu Gln Leu Gln Val Phe Glu Thr Leu Glu Glu Ile Thr Gly Tyr Leu Tyr Ile Ser Ala Trp Pro Asp Ser Leu Pro Asp Leu Ser Val Phe Gln Asn Leu Gln Val Ile Arg Gly Arg Ile Leu His Asn Gly Ala Tyr Ser Leu Thr Leu Gln Gly Leu Gly Ile Ser Trp Leu Gly Leu Arg Ser Leu Arg Glu Leu Gly Ser Gly Leu Ala Leu Ile His His Asn Thr His Leu Cys Phe Val His Thr Val Pro Trp Asp Gln Leu Phe Arg Asn Pro His Gln Ala Leu Leu His Thr Ala Asn Arg Pro Glu Asp Glu Cys Val Gly Glu Gly Leu Ala Cys His Gln Leu Cys Ala Arg Gly His Cys Trp Gly Pro Gly Pro Thr Gln Cys Val Asn Cys Ser Gln Phe Leu Arg Gly Gln Glu Cys Val Glu Glu Cys 

Arg Val Leu Gln Gly Leu Pro Arg Glu Tyr Val Asn Ala Arg His Cys

Leu Pro Cys His Pro Glu Cys Gln Pro Gln Asn Gly Ser Val Thr Cys Phe Gly Pro Glu Ala Asp Gln Cys Val Ala Cys Ala His Tyr Lys Asp Pro Pro Phe Cys Val Ala Arg Cys Pro Ser Gly Val Lys Pro Asp Leu Ser Tyr Met Pro Ile Trp Lys Phe Pro Asp Glu Glu Gly Ala Cys Gln Pro Cys Pro Ile Asn Cys Thr His Ser Cys Val Asp Leu Asp Asp Lys Gly Cys Pro Ala Glu Gln Arg Ala Ser Pro Leu Thr Ser Ile Ile Ser Ala Val Val Gly Ile Leu Leu Val Val Leu Gly Val Val Phe Gly Ile Leu Ile Lys Arg Arg Gln Gln Lys Ile Arg Lys Tyr Thr Met Arg Arg Leu Leu Gln Glu Thr Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Ala Met Pro Asn Gln Ala Gln Met Arg Ile Leu Lys Glu Thr Glu Leu Arg Lys Val Lys Val Leu Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys Gly Ile Trp Ile Pro Asp Gly Glu Asn Val Lys Ile Pro Val Ala Ile Lys Val Leu Arg Glu Asn Thr Ser Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Gly Val Gly Ser Pro Tyr Val Ser Arg Leu Leu Gly Ile Cys Leu Thr Ser Thr Val Gln Leu Val Thr Gln Leu

- Met Pro Tyr Gly Cys Leu Leu Asp His Val Arg Glu Asn Arg Gly Arg 805 810 815
- Leu Gly Ser Gln Asp Leu Leu Asn Trp Cys Met Gln Ile Ala Lys Gly 820 825 830
- Met Ser Tyr Leu Glu Asp Val Arg Leu Val His Arg Asp Leu Ala Ala 835 840 845
- Arg Asn Val Leu Val Lys Ser Pro Asn His Val Lys Ile Thr Asp Phe 850 860
- Gly Leu Ala Arg Leu Leu Asp Ile Asp Glu Thr Glu Tyr His Ala Asp 865 870 875 880
- Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu Glu Ser Ile Leu Arg 885 890 895
- Arg Arg Phe Thr His Gln Ser Asp Val Trp Ser Tyr Gly Val Thr Val 900 905 910
- Trp Glu Leu Met Thr Phe Gly Ala Lys Pro Tyr Asp Gly Ile Pro Ala 915 920 925
- Arg Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg Leu Pro Gln Pro 930 935 940
- Pro Ile Cys Thr Ile Asp Val Tyr Met Ile Met Val Lys Cys Trp Met 945 950 955 960
- Ile Asp Ser Glu Cys Arg Pro Arg Phe Arg Glu Leu Val Ser Glu Phe 965 970 975
- Ser Arg Met Ala Arg Asp Pro Gln Arg Phe Val Val Ile Gln Asn Glu 980 985 990
- Asp Leu Gly Pro Ala Ser Pro Leu Asp Ser Thr Phe Tyr Arg Ser Leu 995 1000 1005
- Leu Glu Asp Asp Asp Met Gly Asp Leu Val Asp Ala Glu Glu Tyr 1010 1015 1020
- Leu Val Pro Gln Gln Gly Phe Phe Cys Pro Asp Pro Ala Pro Gly 1025 1030 1035

Ala Gly Gly Met Val His His Arg His Arg Ser Ser Ser Thr Arg 1040 1045 Ser Gly Gly Gly Asp Leu Thr Leu Gly Leu Glu Pro Ser Glu Glu 1055 1060 Glu Ala Pro Arg Ser Pro Leu Ala Pro Ser Glu Gly Ala Gly Ser 1075 Asp Val Phe Asp Gly Asp Leu Gly Met Gly Ala Ala Lys Gly Leu 1090 Gln Ser Leu Pro Thr His Asp Pro Ser Pro Leu Gln Arg Tyr Ser 1105 1110 Glu Asp Pro Thr Val Pro Leu Pro Ser Glu Thr Asp Gly Tyr Val 1115 1120 Ala Pro Leu Thr Cys Ser Pro Gln Pro Glu Tyr Val Asn Gln Pro 1135 Asp Val Arg Pro Gln Pro Pro Ser Pro Arg Glu Gly Pro Leu Pro 1150 Ala Ala Arg Pro Ala Gly Ala Thr Leu Glu Arg Pro Lys Thr Leu 1160 1165 1170 Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val Phe Ala Phe Gly 1175 1180 1185

Gly Ala Val Glu Asn Pro Glu Tyr Leu Thr Pro Gln Gly Gly Ala 1190 1195 1200

Ala Pro Gln Pro His Pro Pro Pro Ala Phe Ser Pro Ala Phe Asp 1205 1210 1215

Asn Leu Tyr Tyr Trp Asp Gln Asp Pro Pro Glu Arg Gly Ala Pro 1220 1230

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					gac Asp										707
					cct Pro										755
					cca Pro										803
					cag Gln										851
					ggc Gly 230				-					_	899
_	_	_		_	acg Thr			_		_	-	_	_	_	947
					agt Ser										995
					gac Asp										1043
					gcc Ala										1091
					gga Gly 310										1139
					gag Glu										1187
					gta Val										1235
			_		acc Thr	_	_		_			_		_	1283
					agc Ser										1331
					gtt Val 390										1379

					gag Glu										1427
					gac Asp										1475
					cat His										1523
					ctg Leu										1571
	_	_			cac His 470	_				_		_			1619
_			_	_	ctc Leu				_	_	_				1667
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					cgt Arg										1763
					cag Gln										1811
					ggg Gly 550										1859
					ccc Pro										1907
					gct Ala										1955
					gtg Val										2003
					atc Ile										2051
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-	 -	cca Pro	_		_	-	-	_					2147
-		gtg Val 660		-	_	_		_			 -		2195
		atc Ile											2243
		ctg Leu										:	2291
	_	ccc Pro		_	_	_	_			_		 :	2339
		ctg Leu										:	2387
		tgg Trp 740										:	2435
		ttg Leu										:	2483
		gcg Ala										:	2531
		ggc Gly										:	2579
		tat Tyr										2	2627
		tcc Ser 820										2	2675
		tac Tyr										2	2723
		gtg Val										2	2771
		gca Ala										2	2819
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	cgc Arg	Arg												y Vai		2915
	tgg Trp												o Gl			2963
	cgg Arg 930											u Ar				3011
	cca Pro										Met					3059
	att (			_		-	_	_			_				r Glu	3107
	tcc Ser	Arg												e Gli		3155
	Asp					_		Met				nr Pl			cgt tca Arg Ser	3203
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	ctg Leu 1025	Val		_	cag Gln		Ph				ro Ž	gac Asp 1035		gcc Ala		3293
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	agt Ser 1055						Th				eu (	gag Glu 1065				3383
_	gag Glu 1070						Le				er (	gaa Glu 1080				3428
	gat Asp 1085				ggt Gly		Le				ly V			aaa Lys		3473
_	cag Gln 1100	_					Ās			-	ro I	cta Leu 1110	_			3518

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	gct Ala 1130														3608
	gag Glu 1145	-			_			_			gag Glu 1155			_	3653
	ccc Pro 1160										aga Arg 1170				3698
	tct Ser 1175										gtt Val 1185		gcc Ala		3743
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	gcc Ala 1205										agc Ser 1215				3833
	aac Asn 1220												ggt Gly		3878
	cca Pro 1235										gag Glu 1245				3923
	cta Leu 1250					cca Pro 1255		tga	ggto	cacat	gt go	cagad	catco		3970
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Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr 50 55 60

Leu Pro Ala Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val 65 70 75 80

Gln Gly Tyr Met Leu Ile Ala His Asn Arg Val Lys His Val Pro Leu 85 90 95

Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Lys Tyr 100 105 110

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Ala Ala Pro Gly Arg Thr Pro Glu Gly Leu Arg Glu Leu Gln Leu Arg 130 135 140

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Asn Pro Ser Ser Gly Val Ala Pro Leu Lys Pro Glu His Leu Gln Val Phe Glu Thr Leu Glu Glu Ile Thr Gly Tyr Leu Tyr Ile Ser Ala Trp Pro Glu Ser Phe Gln Asp Leu Ser Val Phe Gln Asn Leu Arg Val Ile Arg Gly Arg Ile Leu His Asp Gly Ala Tyr Ser Leu Thr Leu Gln Gly Leu Gly Ile His Ser Leu Gly Leu Arg Ser Leu Arg Glu Leu Gly Ser Gly Leu Ala Leu Ile His Arg Asn Thr His Leu Cys Phe Val Asn Thr Val Pro Trp Asp Gln Leu Phe Arg Asn Pro His Gln Ala Leu Leu His Ser Gly Asn Arg Pro Glu Glu Ala Cys Gly Leu Glu Gly Leu Val Cys Asn Ser Leu Cys Ala Arg Gly His Cys Trp Gly Pro Gly Pro Thr Gln Cys Val Asn Cys Ser Gln Phe Leu Arg Gly Gln Glu Cys Val Glu Glu Cys Arg Val Trp Lys Gly Leu Pro Arg Glu Tyr Val Arg Gly Lys His Cys Leu Pro Cys His Pro Glu Cys Gln Pro Gln Asn Ser Ser Glu Thr Cys Tyr Gly Ser Glu Ala Asp Gln Cys Glu Ala Cys Ala His Tyr Lys Asp Ser Ser Cys Val Ala Arg Cys Pro Ser Gly Val Lys Pro Asp Leu Ser Tyr Met Pro Ile Trp Lys Tyr Pro Asp Glu Glu Gly Ile Cys 

Gln Pro Cys Pro Ile Asn Cys Thr His Ser Cys Val Asp Leu Asp Glu Arg Gly Cys Pro Ala Glu Gln Arg Ala Ser Pro Val Thr Phe Ile Ile Ala Thr Val Val Gly Val Leu Leu Phe Leu Ile Ile Val Val Ile Gly Ile Leu Ile Lys Arg Arg Gln Lys Ile Arg Lys Tyr Thr Met Arg Arg Leu Leu Gln Glu Thr Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Ala Val Pro Asn Gln Ala Gln Met Arg Ile Leu Lys Glu Thr Glu Leu Arg Lys Leu Lys Val Leu Gly Ser Gly Ala Phe Gly Thr Val Tyr Lys Gly Ile Trp Ile Pro Asp Gly Glu Asn Val Lys Ile Pro Val Ala Ile Lys Val Leu Arg Glu Asn Thr Ser Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Gly Val Gly Ser Pro Tyr Val Ser Arg Leu Leu Gly Ile Cys Leu Thr Ser Thr Val Gln Leu Val Thr Gln Leu Met Pro Tyr Gly Cys Leu Leu Asp His Val Arg Glu His Arg Gly Arg Leu Gly Ser Gln Asp Leu Leu Asn Trp Cys Val Gln Ile Ala Lys Gly Met Ser Tyr Leu Glu Glu Val Arg Leu Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys Ser Pro Asn His Val Lys Ile Thr Asp

- Phe Gly Leu Ala Arg Leu Leu Asp Ile Asp Glu Thr Glu Tyr His Ala 865 870 875 880
- Asp Gly Gly Lys Val Pro Ile Lys Trp Met Ala Leu Glu Ser Ile Leu 885 890 895
- Arg Arg Arg Phe Thr His Gln Ser Asp Val Trp Ser Tyr Gly Val Thr 900 905 910
- Val Trp Glu Leu Met Thr Phe Gly Ala Lys Pro Tyr Asp Gly Ile Pro 915 920 925
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- Pro Pro Ile Cys Thr Ile Asp Val Tyr Met Ile Met Val Lys Cys Trp 945 950 955 960
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- Glu Asp Leu Gly Pro Ser Ser Pro Met Asp Ser Thr Phe Tyr Arg Ser 995 1000 1005
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- Gly Thr Gly Ser Thr Ala His Arg Arg His Arg Ser Ser Ser Ala 1040 1045 1050
- Arg Ser Gly Gly Glu Leu Thr Leu Gly Leu Glu Pro Ser Glu 1055 1060 1065
- Glu Glu Pro Pro Arg Ser Pro Leu Ala Pro Ser Glu Gly Ala Gly 1070 1075 1080
- Ser Asp Val Phe Asp Gly Asp Leu Ala Val Gly Val Thr Lys Gly 1085 1090 1095

Leu Gln Ser Leu Ser Pro His Asp Leu Ser Pro Leu Gln Arg Tyr 1100 1105 Ser Glu Asp Pro Thr Leu Pro Leu Pro Pro Glu Thr Asp Gly Tyr 1115 1120 Val Ala Pro Leu Ala Cys Ser Pro Gln Pro Glu Tyr Val Asn Gln 1135 Pro Glu Val Arg Pro Gln Ser Pro Leu Thr Pro Glu Gly Pro Pro 1150 Pro Pro Ile Arg Pro Ala Gly Ala Thr Leu Glu Arg Pro Lys Thr 1165 Leu Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val Phe Ala Phe 1175 1180 Gly Gly Ala Val Glu Asn Pro Glu Tyr Leu Ala Pro Arg Ala Gly 1195 Thr Ala Ser Gln Pro His Pro Ser Pro Ala Phe Ser Pro Ala Phe 1210 1215 Asp Asn Leu Tyr Tyr Trp Asp Gln Asn Ser Ser Glu Gln Gly Pro 1220 1225 1230 Pro Pro Ser Thr Phe Glu Gly Thr Pro Thr Ala Glu Asn Pro Glu 1235 1240 1245 Tyr Leu Gly Leu Asp Val Pro Val 1250 1255 <210> 14 <211> 5511 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (194)..(4222) <400> 14 acacacac acccctcccc tgccatccct ccccqqactc cggctccggc tccgattgca

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cgt cca gtc tct cta of Arg Pro Val Ser Leu 1				
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Leu Tyr Lys Leu Tyr Glu Arg Cys Glu Val Val Met Gly Asn Leu Glu 50 55 60

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Arg Glu Val Thr Gly Tyr Val Leu Val Ala Met Asn Glu Phe Ser Thr 85

Leu Pro Leu Pro Asn Leu Arg Val Val Arg Gly Thr Gln Val Tyr Asp 100

Gly Lys Phe Ala Ile Phe Val Met Leu Asn Tyr Asn Thr Asn Ser Ser 115 120 125

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Gly Gly Val Tyr Ile Glu Lys Asn Asp Lys Leu Cys His Met Asp Thr 145 150 155 160

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Arg Cys Trp Gly Pro Gly Ser Glu Asp Cys Gln Thr Leu Thr Lys Thr 195 200 205

Ile Cys Ala Pro Gln Cys Asn Gly His Cys Phe Gly Pro Asn Pro Asn 210 215 220

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Thr Asp Cys Phe Ala Cys Arg His Phe Asn Asp Ser Gly Ala Cys Val 245 250 255

Pro Arg Cys Pro Gln Pro Leu Val Tyr Asn Lys Leu Thr Phe Gln Leu 260 265 270

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Ser Cys Pro His Asn Phe Val Val Asp Gln Thr Ser Cys Val Arg Ala 290 295 300

Cys Pro Pro Asp Lys Met Glu Val Asp Lys Asn Gly Leu Lys Met Cys 305 310 315 320

Glu Pro Cys Gly Gly Leu Cys Pro Lys Ala Cys Glu Gly Thr Gly Ser 325 330 335

Gly Ser Arg Phe Gln Thr Val Asp Ser Ser Asn Ile Asp Gly Phe Val 340 345 350

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Asn Val Phe Arg Thr Val Arg Glu Ile Thr Gly Tyr Leu Asn Ile Gln 385 390 395 400

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Met Lys Asn Leu Asn Val Thr Ser Leu Gly Phe Arg Ser Leu Lys Glu 435 440 445

Ile Ser Ala Gly Arg Ile Tyr Ile Ser Ala Asn Arg Gln Leu Cys Tyr 450 455 460

His His Ser Leu Asn Trp Thr Lys Val Leu Arg Gly Pro Thr Glu Glu 465 470 475 480

Arg Leu Asp Ile Lys His Asn Arg Pro Arg Arg Asp Cys Val Ala Glu 485 490 495

Gly Lys Val Cys Asp Pro Leu Cys Ser Ser Gly Gly Cys Trp Gly Pro 500 505 510

Gly Pro Gly Gln Cys Leu Ser Cys Arg Asn Tyr Ser Arg Gly Gly Val 515 520 525

Cys Val Thr His Cys Asn Phe Leu Asn Gly Glu Pro Arg Glu Phe Ala 530 535 540

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- Val Leu Gly Ala Lys Gly Pro Ile Tyr Lys Tyr Pro Asp Val Gln Asn 595 600 605
- Glu Cys Arg Pro Cys His Glu Asn Cys Thr Gln Gly Cys Lys Gly Pro 610 615 620
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- Ala His Ile Val Arg Leu Leu Gly Leu Cys Pro Gly Ser Ser Leu Gln
  770 780
- Leu Val Thr Gln Tyr Leu Pro Leu Gly Ser Leu Leu Asp His Val Arg 785 790 795 800

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- Gln Ile Ala Lys Gly Met Tyr Tyr Leu Glu Glu His Gly Met Val His 820  $\,$  825  $\,$  830  $\,$
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	g tac u Tyr 50											t Gly				192
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						aag Lys										2160
						att Ile										2208
gtc Val	tgc C <b>y</b> s	att Ile	aaa Lys 740	gtc Val	atc Ile	gag Glu	gac Asp	aag Lys 745	agt Ser	Gly ggg	cgg Arg	cag Gln	agt Ser 750	ttt Phe	cag Gln	2256
						ctg Leu										2304
						ctg Leu 775										2352
						ggc Gly										2400
cgt Arg	gag Glu	aca Thr	ctg Leu	gga Gly 805	cca Pro	cag Gln	ctg Leu	ctg Leu	ctc Leu 810	aac Asn	tgg Trp	gga Gly	gta Val	caa Gln 815	att Ile	2448

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	gcg Ala					_		_			_		_	_		2544
	gat Asp 850						_	_	_		_	_	-	_		2592
	cac His															2640
	cac His															2688
	acc Thr															2736
	cga Arg	_	_	_			_	_	_		_					2784
_	cag Gln 930		_		_			_	_		_	_	_	-	_	2832
	tgg Trp															2880
	gag Glu															2928
	aga Arg															2976
ctg Leu	agc Ser	acc Thr 995	aaa Lys	gag Glu	ttg Leu	cag Gln	gat Asp 1000	Ala	ı gaçı Glu	r ctg . Leu	g gaç n Glu	g cca Pro 100	) As	ac ct sp Le	g gac eu Asp	3024
	gac Asp 1010	Leu			g gag . Glu		. G1				y L∈		gcg a Ala T			3069
	ggt Gly 1025	Ser			agc Ser		Pr				r Le		cc chr A			3114
-	ggg Gly 1040	Ser			ctt Leu		. Se				r Gl		ac a 'yr M	-		3159

						ggg Gly 1060						Ser			3204
						ttc Phe 1075									3249
						acg Thr 1090									3294
						ctc Leu 1105									3339
						agc Ser 1120									3384
	cat His 1130					agc Ser 1135							_	ctc Leu	3429
	cca Pro 1145					gaa Glu 1150									3474
	gat Asp 1160	acg Thr	cac His	ctc Leu	aga Arg	ggt Gly 1165	aca Thr	tcc Ser	tct Ser	tcc Ser	cgg Arg 1170	gaa Glu	ggc Gly		3519
	tcg Ser 1175	tca Ser	gta Val	ggt Gly	ctc Leu	agt Ser 1180	tct Ser	gtg Val	ctg Leu	ggt Gly	acc Thr 1185	gaa Glu	gag Glu		3564
						gaa Glu 1195									3609
	ccg Pro 1205					aga Arg 1210									3654
						ggt Gly 1225									3699
						ctc Leu 1240									3744
-	ggc Gly 1250					gag Glu 1255									3789
cgt Arg	ggt Gly 1265	gcg Ala	ggc Gly	ggt Gly	tcc Ser	gga Gly 1270	ggg Gly	gat Asp	tat Tyr	gca Ala	gct Ala 1275	atg Met	ggg Gly	gcc Ala	3834

tgc cc Cys Pro	o Ála	_	_	ggg Gly 1285		_	_	-	_		-	3879
ggg cc Gly Pro 12	o Gly			ccc Pro 1300								3924
act cte Thr Let	ı Arg											3969
gat tac Asp Ty: 13	r Trp			ctt Leu 1330		_	_		_	_	_	4014
att tga Ile	а.											4020

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Leu Asn Gly Leu Ser Val Thr Gly Asp Ala Asp Asn Gln Tyr Gln Thr 35 40 45

Leu Tyr Lys Leu Tyr Glu Lys Cys Glu Val Val Met Gly Asn Leu Glu 50 55 60

Ile Val Leu Thr Gly His Asn Ala Asp Leu Ser Phe Leu Gln Trp Ile 70 75 80

Arg Glu Val Thr Gly Tyr Val Leu Val Ala Met Asn Glu Phe Ser Val 85 90 95

Leu Pro Leu Pro Asn Leu Arg Val Val Arg Gly Thr Gln Val Tyr Asp 100 105 110

Gly Lys Phe Ala Ile Phe Val Met Leu Asn Tyr Asn Thr Asn Ser Ser 115 120 125

His Ala Leu Arg Gln Leu Arg Phe Thr Gln Leu Thr Glu Ile Leu Leu Gly Gly Val Tyr Ile Glu Lys Asn Asp Lys Leu Cys His Met Asp Thr Ile Asp Trp Arg Asp Ile Val Arg Val Pro Asp Ala Glu Ile Val Val Lys Asn Asn Gly Gly Asn Cys Pro Pro Cys His Glu Val Cys Lys Gly Arg Cys Trp Gly Pro Gly Pro Glu Asp Cys Gln Ile Leu Thr Lys Thr Ile Cys Ala Pro Gln Cys Asn Gly Arg Cys Phe Gly Pro Asn Pro Asn Gln Cys Cys His Asp Glu Cys Ala Gly Gly Cys Ser Gly Pro Gln Asp Thr Asp Cys Phe Ala Cys Arg His Phe Asn Asp Ser Gly Ala Cys Val Pro Arg Cys Pro Ala Pro Leu Val Tyr Asn Lys Leu Thr Phe Gln Leu Glu Pro Asn Pro His Ile Lys Tyr Gln Tyr Gly Gly Val Cys Val Ala Ser Cys Pro His Asn Phe Val Val Asp Gln Thr Phe Cys Val Arg Ala Cys Pro Ala Asp Lys Met Glu Val Asp Lys Asn Gly Leu Lys Met Cys Glu Pro Cys Arg Gly Leu Cys Pro Lys Ala Cys Glu Gly Thr Gly Ser Gly Ser Arg Tyr Gln Thr Val Asp Ser Ser Asn Ile Asp Gly Phe Val Asn Cys Thr Lys Ile Leu Gly Asn Leu Asp Phe Leu Ile Thr Gly Leu

Asn Gly Asp Pro Trp His Lys Ile Pro Ala Leu Asp Pro Glu Lys Leu 370 375 Asn Val Phe Arg Thr Val Arg Glu Ile Thr Gly Tyr Leu Asn Ile Gln Ser Trp Pro Pro His Met His Asn Phe Ser Val Phe Ser Asn Leu Thr 410 Thr Ile Gly Gly Arg Ser Leu Tyr Asn Arg Gly Phe Ser Leu Leu Ile 425 Met Lys Asn Leu Asn Val Thr Ser Leu Gly Phe Arg Ser Leu Lys Glu 440 Ile Ser Ala Gly Arg Val Tyr Ile Ser Ala Asn Gln Gln Leu Cys Tyr His His Ser Leu Asn Trp Thr Arg Leu Leu Arg Gly Pro Ala Glu Glu Arg Leu Asp Ile Lys Tyr Asn Arg Pro Leu Gly Glu Cys Val Ala Glu 485 490 Gly Lys Val Cys Asp Pro Leu Cys Ser Ser Gly Gly Cys Trp Gly Pro 500 505 510 Gly Pro Gly Gln Cys Leu Ser Cys Arg Asn Tyr Ser Arg Glu Gly Val 515 520 Cys Val Thr His Cys Asn Val Leu Gln Gly Glu Pro Arg Glu Phe Val 530 535

His Glu Ala His Cys Phe Ser Cys His Pro Glu Cys Gln Pro Met Glu545550555560

Gly Thr Ser Thr Cys Asn Gly Ser Gly Ser Asp Ala Cys Ala Arg Cys 565 570 575

Ala His Phe Arg Asp Gly Pro His Cys Val Asn Ser Cys Pro His Gly 580 585 590

Ile Leu Gly Ala Lys Gly Pro Ile Tyr Lys Tyr Pro Asp Ala Gln Asn 595 600 605

Glu Cys Arg Pro Cys His Glu Asn Cys Thr Gln Gly Cys Lys Gly Pro 610 615 620

Glu Leu Gln Asp Cys Leu Gly Gln Ala Glu Val Leu Met Ser Lys Pro 625 630 635 640

His Leu Val Ile Ala Val Thr Val Gly Leu Thr Val Ile Phe Leu Ile 645 650 655

Leu Gly Gly Ser Phe Leu Tyr Trp Arg Gly Arg Arg Ile Gln Asn Lys 660 665 670

Arg Ala Met Arg Arg Tyr Leu Glu Arg Gly Glu Ser Ile Glu Pro Leu 675 680 685

Asp Pro Ser Glu Lys Ala Asn Lys Val Leu Ala Arg Ile Phe Lys Glu 690 695 700

Thr Glu Leu Arg Lys Leu Lys Val Leu Gly Ser Gly Val Phe Gly Thr 705 710 715 720

Val His Lys Gly Ile Trp Ile Pro Glu Gly Glu Ser Ile Lys Ile Pro 725 730 735

Val Cys Ile Lys Val Ile Glu Asp Lys Ser Gly Arg Gln Ser Phe Gln 740 745 750

Ala Val Thr Asp His Met Leu Ala Val Gly Ser Leu Asp His Ala His 755 760 765

Ile Val Arg Leu Leu Gly Leu Cys Pro Gly Ser Ser Leu Gln Leu Val 770 775 780

Thr Gln Tyr Leu Pro Leu Gly Ser Leu Leu Asp His Val Arg Gln His 785 790 795 800

Arg Glu Thr Leu Gly Pro Gln Leu Leu Asn Trp Gly Val Gln Ile 805 810 815

Ala Lys Gly Met Tyr Tyr Leu Glu Glu His Ser Met Val His Arg Asp 820 825 830

Leu Ala Leu Arg Asn Val Met Leu Lys Ser Pro Ser Gln Val Gln Val 835 840 845

- Ala Asp Phe Gly Val Ala Asp Leu Leu Pro Pro Asp Asp Lys Gln Leu 850 855 860
- Leu His Ser Glu Ala Lys Thr Pro Ile Lys Trp Met Ala Leu Glu Ser 865 870 875 880
- Ile His Phe Gly Lys Tyr Thr His Gln Ser Asp Val Trp Ser Tyr Gly 885 890 895
- Val Thr Val Trp Glu Leu Met Thr Phe Gly Ala Glu Pro Tyr Ala Gly 900 905 910
- Leu Arg Leu Ala Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg Leu 915 920 925
- Ala Gln Pro Gln Ile Cys Thr Ile Asp Val Tyr Met Val Met Val Lys 930 935 940
- Cys Trp Met Ile Asp Glu Asn Ile Arg Pro Thr Phe Lys Glu Leu Ala 945 950 955 960
- Asn Glu Phe Thr Arg Met Ala Arg Asp Pro Pro Arg Tyr Leu Val Ile 965 970 975
- Lys Arg Ala Ser Gly Pro Gly Ile Pro Pro Ala Ala Glu Pro Ser Ala 980 985 990
- Leu Ser Thr Lys Glu Leu Gln Asp Ala Glu Leu Glu Pro Asp Leu Asp 995 1000 1005
- Leu Asp Leu Asp Val Glu Val Glu Glu Glu Gly Leu Ala Thr Thr 1010 1015 1020
- Leu Gly Ser Ala Leu Ser Leu Pro Thr Gly Thr Leu Thr Arg Pro 1025 1030 1035
- Arg Gly Ser Gln Ser Leu Leu Ser Pro Ser Ser Gly Tyr Met Pro 1040 1045 1050
- Met Asn Gln Ser Asn Leu Gly Glu Ala Cys Leu Asp Ser Ala Val 1055 1060 1065
- Leu Gly Gly Arg Glu Gln Phe Ser Arg Pro Ile Ser Leu His Pro 1070 1075 1080

- Ile Pro Arg Gly Arg Gln Thr Ser Glu Ser Ser Glu Gly His Val 1085 1090 1095
- Thr Gly Ser Glu Ala Glu Leu Gln Glu Arg Val Ser Met Cys Arg 1100 \$1105\$ \$1110
- Ser Arg Ser Arg Ser Pro Arg Pro Arg Gly Asp Ser Ala 1115 1120 1125
- Tyr His Ser Gln Arg His Ser Leu Leu Thr Pro Val Thr Pro Leu 1130 1135 1140
- Ser Pro Pro Gly Leu Glu Glu Glu Asp Gly Asn Gly Tyr Val Met 1145 1150 1155
- Pro Asp Thr His Leu Arg Gly Thr Ser Ser Ser Arg Glu Gly Thr 1160 1165 1170
- Leu Ser Ser Val Gly Leu Ser Ser Val Leu Gly Thr Glu Glu Glu 1175 1180 1185
- Asp Glu Asp Glu Glu Tyr Glu Tyr Met Asn Arg Lys Arg Arg Gly 1190 1195 1200
- Ser Pro Ala Arg Pro Pro Arg Pro Gly Ser Leu Glu Glu Leu Gly 1205 1210 1215
- Tyr Glu Tyr Met Asp Val Gly Ser Asp Leu Ser Ala Ser Leu Gly 1220 1225 1230
- Ser Thr Gln Ser Cys Pro Leu His Pro Met Ala Ile Val Pro Ser 1235 1240 1245
- Ala Gly Thr Thr Pro Asp Glu Asp Tyr Glu Tyr Met Asn Arg Arg 1250 1255 1260
- Arg Gly Ala Gly Gly Ser Gly Gly Asp Tyr Ala Ala Met Gly Ala 1265 1270 1275
- Cys Pro Ala Ala Glu Gln Gly Tyr Glu Glu Met Arg Ala Phe Gln 1280 1285 1290
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Thr Leu Arg Ser Leu Glu Ala Thr Asp Ser Ala Phe Asp Asn Pro

1315

		gaa Glu								486
		tat Tyr								534
		tcc Ser	_						_	582
		tgc Cys								630
		tgc Cys 205								678
		tgc Cys								726
		ggc Gly		-	_	_	_	_		774
		aat Asn								822
		aat Asn				_				870
		tat Tyr 285								918
		tcc Ser								966
		gaa Glu								1014
		gct Ala								1062
		gat Asp								1110
		aat Asn 365								1158

_			_	att Ile	_	_		_				_		_	٠	1206
				gag Glu												1254
				gac Asp												1302
				tat Tyr												1350
				cta Leu 445												1398
				act Thr												1446
				ctc Leu		-				_	_		_			1494
				gct Ala	_				-	-		_		-		1542
				agt Ser												1590
				cgc Arg 525												1638
				ggt Gly												1686
				ccc Pro												1734
	_		_	ggt Gly		_		_		-	_					1782
				tgt Cys												1830
				ttc Phe 605			-	-		_			_			1878

			tgc Cys											1926
_	_		cca Pro		_									1974
			att Ile											2022
			ctg Leu											2070
			gcc Ala 685											2118
			agt Ser							_			_	2166
			gag Glu	_	_		_		_					2214
			tat Tyr					_		_		-		2262
			 gct Ala		_									2310
_	_		 ttc Phe 765	_	_	_	_	_		_	-	_	_	2358
			gtc Val											2406
			caa Gln											2454
			gat Asp											2502
			aag Lys											2550
			gca Ala 845											2598

							Ar		c ttg u Leu			Asp	2646
				_	 	Lys		-	a att o Ile			_	2694
									t cag s Gln 900	Ser			2742
									c ttt r Phe 5				2790
								o As	t tta p Leu				2838
									t gac e Asp				2886
									t aga r Arg				2934
									a gac g Asp 980				2982
									t ccc u Pro 5				3030
gac Asp 1000	Ser	aag Lys		ı As			sp (		gag Glu		_		3075
gat Asp 1015	Met			Gl			al :		cag Gln				3120
atc Ile 1030	Pro			Th			la <i>i</i>		att (				3165
agg Arg 1045	Ser			Se			ro Z		tac a				3210
tca Ser 1060	Gly			Ту			Ly (		ttt o				3255
caa Gln 1075	Gly	gtg Val		Ту			co :		agc a				3300

gaa Glu 1090	Ala		cag Gln 1095	Gly				Ile				3	345
tcc Ser 1105	Cys		acc Thr 1110									3	390
caa Gln 1120	Glu		acc Thr 1125	Gln				Asp				3	435
ttt Phe 1135	Ala		agc Ser 1140									3	480
tac Tyr 1150	Met		cga Arg 1155									3	525
cca Pro 1165			cct Pro 1170									3	570
ctt Leu 1180			aat Asn 1185									3	61 <sup>.</sup> 5
cca Pro 1195			gat Asp 1200						-	tac Tyr		3	660
aac Asn 1210			acc Thr 1215									3	705
aac Asn 1225			cca Pro 1230									3	750
cct Pro 1240			cac His 1245									3	795
cac His 1255			cag Gln 1260									3	840
cag Gln 1270		 	cgg Arg 1275			_				_		3	885
ctc Leu 1285			ctg Leu 1290									3	930
cct Pro 1300			aat Asn 1305			taa	gctca	agtto	jt gg	jtttt	ttag	39	980

4040 gtggagagac acacctgctc caatttcccc accccctct ctttctctgg tggtcttcct 4100 tctaccccaa ggccagtagt tttgacactt cccagtggaa gatacagaga tgcaatgata gttatgtgct tacctaactt gaacattaga gggaaagact gaaagagaaa gataggagga 4160 accacaatgt ttcttcattt ctctgcatgg gttggtcagg agaatgaaac agctagagaa 4220 ggaccagaaa atgtaaggca atgctgccta ctatcaaact agctgtcact ttttttcttt 4280 4340 ggttgaaaca cccatgctat ctgttcctat ctgcaggaac tgatgtgtgc atatttagca 4400 4460 tccctggaaa tcataataaa gtttccatta gaacaaaaga ataacatttt ctataacata tgatagtgtc tgaaattgag aatccagttt ctttccccag cagtttctgt cctagcaagt 4520 aagaatggcc aactcaactt tcataattta aaaatctcca ttaaagttat aactagtaat 4580 4640 tttgctcccc tatttttggc tttaatttct aattgcaaag atgtttacat caaagcttct 4700 tcacagaatt taagcaagaa atattttaat atagtgaaat ggccactact ttaagtatac 4760 aatctttaaa ataagaaagg gaggctaata tttttcatgc tatcaaatta tcttcaccct 4820 catcctttac atttttcaac atttttttt ctccataaat gacactactt gataggccgt 4880 tggttgtctg aagagtagaa gggaaactaa gagacagttc tctgtggttc aggaaaacta 4940 ctgatacttt caggggtggc ccaatgaggg aatccattga actggaagaa acacactgga 5000 5060 ttgggtatgt ctacctggca gatactcaga aatgtagttt gcacttaagc tgtaatttta 5120 tttgttcttt ttctgaactc cattttggat tttgaatcaa gcaatatgga agcaaccagc aaattaacta atttaagtac atttttaaaa aaagagctaa gataaagact gtggaaatgc 5180 caaaccaagc aaattaggaa ccttgcaacg gtatccaggg actatgatga gaggccagca 5240 5300 cattatette atatgteace tttgetacge aaggaaattt gtteagtteg tataettegt aagaaggaat gcgagtaagg attggcttga attccatgga atttctagta tgagactatt 5360 tatatgaagt agaaggtaac tctttgcaca taaattggta taataaaaag aaaaacacaa 5420 acattcaaag cttagggata ggtccttggg tcaaaagttg taaataaatg tgaaacatct 5480 5484 tctc

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<sup>&</sup>lt;211> 1308

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

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Glu Asn Lys Leu Ser Ser Leu Ser Asp Leu Glu Gln Gln Tyr Arg Ala 35 40 45

Leu Arg Lys Tyr Tyr Glu Asn Cys Glu Val Val Met Gly Asn Leu Glu 50 55 60

Ile Thr Ser Ile Glu His Asn Arg Asp Leu Ser Phe Leu Arg Ser Val 65 70 75 80

Arg Glu Val Thr Gly Tyr Val Leu Val Ala Leu Asn Gln Phe Arg Tyr 85 90 95

Leu Pro Leu Glu Asn Leu Arg Ile Ile Arg Gly Thr Lys Leu Tyr Glu 100 105 110

Asp Arg Tyr Ala Leu Ala Ile Phe Leu Asn Tyr Arg Lys Asp Gly Asn 115 120 125

Phe Gly Leu Gl<br/>n Glu Leu Gly Leu Lys As<br/>n Leu Thr Glu Ile Leu As<br/>n 130 135 140

Gly Gly Val Tyr Val Asp Gln Asn Lys Phe Leu Cys Tyr Ala Asp Thr 145 150 155 160

Ile His Trp Gln Asp Ile Val Arg Asn Pro Trp Pro Ser Asn Leu Thr 165 170 175

Leu Val Ser Thr Asn Gly Ser Ser Gly Cys Gly Arg Cys His Lys Ser 180 185 190

Cys Thr Gly Arg Cys Trp Gly Pro Thr Glu Asn His Cys Gln Thr Leu 195 200 205

Thr Arg Thr Val Cys Ala Glu Gln Cys Asp Gly Arg Cys Tyr Gly Pro 210 215 220

Tyr Val Ser Asp Cys Cys His Arg Glu Cys Ala Gly Gly Cys Ser Gly 235 Pro Lys Asp Thr Asp Cys Phe Ala Cys Met Asn Phe Asn Asp Ser Gly 245 250 Ala Cys Val Thr Gln Cys Pro Gln Thr Phe Val Tyr Asn Pro Thr Thr 265 Phe Gln Leu Glu His Asn Phe Asn Ala Lys Tyr Thr Tyr Gly Ala Phe Cys Val Lys Lys Cys Pro His Asn Phe Val Val Asp Ser Ser Ser Cys 290 295 Val Arg Ala Cys Pro Ser Ser Lys Met Glu Val Glu Glu Asn Gly Ile 310 Lys Met Cys Lys Pro Cys Thr Asp Ile Cys Pro Lys Ala Cys Asp Gly 330 335 Ile Gly Thr Gly Ser Leu Met Ser Ala Gln Thr Val Asp Ser Ser Asn 340 345 350 Ile Asp Lys Phe Ile Asn Cys Thr Lys Ile Asn Gly Asn Leu Ile Phe 355 360 365 Leu Val Thr Gly Ile His Gly Asp Pro Tyr Asn Ala Ile Glu Ala Ile 370 375 Asp Pro Glu Lys Leu Asn Val Phe Arg Thr Val Arg Glu Ile Thr Gly 385 390 395 Phe Leu Asn Ile Gln Ser Trp Pro Pro Asn Met Thr Asp Phe Ser Val 405 Phe Ser Asn Leu Val Thr Ile Gly Gly Arg Val Leu Tyr Ser Gly Leu Ser Leu Leu Ile Leu Lys Gln Gln Gly Ile Thr Ser Leu Gln Phe Gln Ser Leu Lys Glu Ile Ser Ala Gly Asn Ile Tyr Ile Thr Asp Asn Ser 455

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Asn Leu Cys Tyr Tyr His Thr Ile Asn Trp Thr Thr Leu Phe Ser Thr Ile Asn Gln Arg Ile Val Ile Arg Asp Asn Arg Lys Ala Glu Asn Cys Thr Ala Glu Gly Met Val Cys Asn His Leu Cys Ser Ser Asp Gly Cys Trp Gly Pro Gly Pro Asp Gln Cys Leu Ser Cys Arg Arg Phe Ser Arg Gly Arg Ile Cys Ile Glu Ser Cys Asn Leu Tyr Asp Gly Glu Phe Arg Glu Phe Glu Asn Gly Ser Ile Cys Val Glu Cys Asp Pro Gln Cys Glu Lys Met Glu Asp Gly Leu Leu Thr Cys His Gly Pro Gly Pro Asp Asn Cys Thr Lys Cys Ser His Phe Lys Asp Gly Pro Asn Cys Val Glu Lys Cys Pro Asp Gly Leu Gln Gly Ala Asn Ser Phe Ile Phe Lys Tyr Ala Asp Pro Asp Arg Glu Cys His Pro Cys His Pro Asn Cys Thr Gln Gly Cys Asn Gly Pro Thr Ser His Asp Cys Ile Tyr Tyr Pro Trp Thr Gly His Ser Thr Leu Pro Gln His Ala Arg Thr Pro Leu Ile Ala Ala Gly Val Ile Gly Gly Leu Phe Ile Leu Val Ile Val Gly Leu Thr Phe Ala Val Tyr Val Arg Arg Lys Ser Ile Lys Lys Arg Ala Leu Arg Arg Phe Leu Glu Thr Glu Leu Val Glu Pro Leu Thr Pro Ser Gly Thr Ala 

Pro 705	Asn	Gln	Ala	Gln	Leu 710	Arg	Ile	Leu	Lys	Glu 715	Thr	Glu	Leu	Lys	Arg 720
Val	Lys	Val	Leu	Gly 725	Ser	Gly	Ala	Phe	Gly 730	Thr	Val	Tyr	Lys	Gly 735	Ile
Trp	Val	Pro	Glu 740	Gly	Glu	Thr	Val	Lys 745	Ile	Pro	Val	Ala	Ile 750	Lys	Ile
Leu	Asn	Glu 755	Thr	Thr	Gly	Pro	Lys 760	Ala	Asn	Val	Glu	Phe 765	Met	Asp	Glu
Ala	Leu 770	Ile	Met	Ala	Ser	Met 775	Asp	His	Pro	His	Leu 780	Val	Arg	Leu	Leu
Gly 785	Val	Cys	Leu	Ser	Pro 790	Thr	Ile	Gln	Leu	Val 795	Thr	Gln	Leu	Met	Pro 800
His	Gly	Cys	Leu	Leu 805	Glu	Tyr	Val	His	Glu 810	His	Lys	Asp	Asn	Ile 815	Gly
Ser	Gln	Leu	Leu 820	Leu	Asn	Trp	Cys	Val 825	Gln	Ile	Ala	Lys	Gly 830	Met	Met
Tyr	Leu	Glu 835	Glu	Arg	Arg	Leu	Val 840	His	Arg	Asp	Leu	Ala 845	Ala	Arg	Asn
Val	Leu 850	Val	Lys	Ser	Pro	Asn 855	His	Val	Lys	Ile	Thr 860	Asp	Phe	Gly	Leu
Ala 865	Arg	Leu	Leu	Glu	Gly 870	Asp	Glu	Lys	Glu	Tyr 875	Asn	Ala	Asp	Gly	Gly 880
Lys	Met	Pro	Ile	Lys 885	Trp	Met	Ala	Leu	Glu 890	Cys	Ile	His	Tyr	Arg 895	Lys
Phe	Thr	His	Gln 900	Ser	Asp	Val	Trp	Ser 905	Tyr	Gly	Val	Thr	Ile 910	Trp	Glu
Leu	Met	Thr 915	Phe	Gly	Gly	Lys	Pro 920	Tyr	Asp	Gly	Ile	Pro 925	Thr	Arg	Glu
Ile	Pro 930	Asp	Leu	Leu	Glu	Lys 935	Gly	Glu	Arg	Leu	Pro 940	Gln	Pro	Pro	Ile

Cys Thr Ile Asp Val Tyr Met Val Met Val Lys Cys Trp Met Ile Asp 945 950 955 960

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- Ala Asp Ser Arg Pro Lys Phe Lys Glu Leu Ala Ala Glu Phe Ser Arg 965 970 975
- Met Ala Arg Asp Pro Gln Arg Tyr Leu Val Ile Gln Gly Asp Asp Arg 980 985 990
- Met Lys Leu Pro Ser Pro Asn Asp Ser Lys Phe Phe Gln Asn Leu Leu 995 1000 1005
- Asp Glu Glu Asp Leu Glu Asp Met Met Asp Ala Glu Glu Tyr Leu 1010 1015 1020
- Val Pro Gln Ala Phe Asn Ile Pro Pro Pro Ile Tyr Thr Ser Arg 1025 1030 1035
- Ala Arg Ile Asp Ser Asn Arg Ser Glu Ile Gly His Ser Pro Pro 1040 1045 1050
- Pro Ala Tyr Thr Pro Met Ser Gly Asn Gln Phe Val Tyr Arg Asp 1055 1060 1065
- Gly Gly Phe Ala Ala Glu Gln Gly Val Ser Val Pro Tyr Arg Ala 1070 1075 1080
- Pro Thr Ser Thr Ile Pro Glu Ala Pro Val Ala Gln Gly Ala Thr 1085 1090 1095
- Ala Glu Ile Phe Asp Asp Ser Cys Cys Asn Gly Thr Leu Arg Lys 1100 1105 1110
- Pro Val Ala Pro His Val Gln Glu Asp Ser Ser Thr Gln Arg Tyr 1115 1120 1125
- Ser Ala Asp Pro Thr Val Phe Ala Pro Glu Arg Ser Pro Arg Gly 1130 1135 1140
- Glu Leu Asp Glu Glu Gly Tyr Met Thr Pro Met Arg Asp Lys Pro 1145 1150 1155
- Lys Gln Glu Tyr Leu Asn Pro Val Glu Glu Asn Pro Phe Val Ser 1160 1165 1170

Arg Arg Lys Asn Gly Asp Leu Gln Ala Leu Asp Asn Pro Glu Tyr 1175 1180 1185 His Asn Ala Ser Asn Gly Pro Pro Lys Ala Glu Asp Glu Tyr Val 1195 Asn Glu Pro Leu Tyr Leu Asn Thr Phe Ala Asn Thr Leu Gly Lys 1205 Ala Glu Tyr Leu Lys Asn Asn Ile Leu Ser Met Pro Glu Lys Ala 1225 Lys Lys Ala Phe Asp Asn Pro Asp Tyr Trp Asn His Ser Leu Pro 1240 Pro Arg Ser Thr Leu Gln His Pro Asp Tyr Leu Gln Glu Tyr Ser 1250 1255 Thr Lys Tyr Phe Tyr Lys Gln Asn Gly Arg Ile Arg Pro Ile Val 1270 Ala Glu Asn Pro Glu Tyr Leu Ser Glu Phe Ser Leu Lys Pro Gly 1285 Thr Val Leu Pro Pro Pro Tyr Arg His Arg Asn Thr Val Val 1295 1300 1305 <210> 20 <211> 3897 <212> DNA <213> Mus musculus <220> <221> CDS <222> (1)..(3897) <400> 20 atg gca ata cca tca aga cgc aag gac tat cca aaa cta tct gaa atc 48 Met Ala Ile Pro Ser Arg Arg Lys Asp Tyr Pro Lys Leu Ser Glu Ile aca gtg tgc gca gga aca gag aac aaa ctg agc tct ctc tct gac ctg 96 Thr Val Cys Ala Gly Thr Glu Asn Lys Leu Ser Ser Leu Ser Asp Leu 25 gaa cag cag tac cga gcc ttg cgc aaa tac tat gaa aac tgc gag gta 144 Glu Gln Gln Tyr Arg Ala Leu Arg Lys Tyr Tyr Glu Asn Cys Glu Val 35 40

	ggc Gly								192
	ctg Leu								240
	cag Gln								288
	aaa Lys								336
	aaa Lys 115								384
	gaa Glu								432
	tat Tyr								480
	tcc Ser								528
	tgc Cys								576
	tgc Cys 195								624
	tgc Cys								672
	ggc Gly								720
	aat Asn								768
	aat Asn								816
	tat Tyr 275		Cys						864

		tgt Cys						912
		att Ile 310						960
		gga Gly						1008
		aac Asn						1056
		ttt Phe	-			 _		1104
		ata Ile						1152
		ggt Gly 390						1200
		gtt Val						1248
		ctc Leu						1296
		cag Gln			_	 		1344
		agc Ser						1392
		acc Thr 470						1440
		tgt Cys						1488
		tgt Cys						1536
		agg Arg						1584

				gaa Glu					1632
				gaa Glu					1680
				aag Lys					1728
				gat Asp 585					1776
				gat Asp					1824
				ggt Gly					1872
				act Thr					1920
				gga Gly					1968
				gtc Val 665					2016
				gag Glu					2064
Thr				caa Gln					2112
				gtc Val					2160
				cct Pro				:	2208
				gaa Glu 745				;	2256
				atc Ile				:	2304

			ttg Leu										2352
			ccg Pro 790										2400
			gga Gly										2448
			atg Met										2496
			aat Asn								_		2544
			ctg Leu										2592
			ggc Gly 870										2640
			aaa Lys										2688
			gaa Glu										2736
			gaa Glu										2784
			atc Ile										2832
_	 _		gat Asp 950	_	_	-	-				-	_	2880
_			aga Arg	_	_	_	_		_			Val	2928
_	 -	-	cgt Arg	_	_			_		_	_		2976
			ttg Leu			Glu			gac Asp 100	Me	-	g gat t Asp	3024

_		_		_	_	ccc Pro 1015	_	_				Pro			3069
	tac Tyr 1025			_	Thr	aga Arg 1030		_				_	_		3114
						gcc Ala 1045									3159
	gtg Val 1055			_		ggc Gly 1060		-					_		3204
_	ccc Pro 1070		_	_		acc Thr 1075	_					_		-	3249
_	_		_	_	-	gag Glu 1090	_		-	_	tcc Ser 1095	_	_		3294
	acc Thr 1100					gtg Val 1105									3339
_	act Thr 1115					gct Ala 1120									3384
						ctg Leu 1135							act Thr		3429
						caa Gln 1150			-				_		3474
-	cct Pro 1160					agg Arg 1165									3519
gat Asp	aat Asn 1175					agt Ser 1180									3564
_	gat Asp 1190					gag Glu 1195							ttc Phe	_	3609
	gcc Ala 1205	_		_	_	gag Glu 1210		_			_	_	_		3654
						aaa Lys 1225									3699

aac cac age ctg cca ccc cgg age ace ctt cag of Asn His Ser Leu Pro Pro Arg Ser Thr Leu Gln H	
ctg cag gaa tac agc aca aaa tat ttt tat aaa c Leu Gln Glu Tyr Ser Thr Lys Tyr Phe Tyr Lys ( 1250 1255	
atc cgc ccc att gtg gca gag aat cct gag tac c Ile Arg Pro Ile Val Ala Glu Asn Pro Glu Tyr I 1265 1270	
tcg ctg aag cct ggc act atg ctg ccc cct ccg c Ser Leu Lys Pro Gly Thr Met Leu Pro Pro Pro I 1280	
cgg aat act gtg gtg tga Arg Asn Thr Val Val 1295	3897
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Thr Val Cys Ala Gly Thr Glu Asn Lys Leu Ser Ser 20 25	r Leu Ser Asp Leu 30
Glu Gln Gln Tyr Arg Ala Leu Arg Lys Tyr Tyr Glu 35 40	ı Asn Cys Glu Val 45
Val Met Gly Asn Leu Glu Ile Thr Ser Ile Glu His 50 55 60	s Asn Arg Asp Leu
Ser Phe Leu Arg Ser Ile Arg Glu Val Thr Gly Tyr 65 70 75	val Leu Val Ala 80
Leu Asn Gln Phe Arg Tyr Leu Pro Leu Glu Asn Leu 85 90	a Arg Ile Ile Arg 95
Gly Thr Lys Leu Tyr Glu Asp Arg Tyr Ala Leu Ala	a Ile Phe Leu Asn 110
Tyr Arg Lys Asp Gly Asn Phe Gly Leu Gln Glu Leu 115 120	ı Gly Leu Lys Asn 125

Leu Thr Glu Ile Leu Asn Gly Gly Val Tyr Val Asp Gln Asn Lys Phe Leu Cys Tyr Ala Asp Thr Ile His Trp Gln Asp Ile Val Arg Asn Pro Trp Pro Ser Asn Met Thr Leu Val Ser Thr Asn Gly Ser Ser Gly Cys Gly Arg Cys His Lys Ser Cys Thr Gly Arg Cys Trp Gly Pro Thr Glu Asn His Cys Gln Thr Leu Thr Arg Thr Val Cys Ala Glu Gln Cys Asp Gly Arg Cys Tyr Gly Pro Tyr Val Ser Asp Cys Cys His Arg Glu Cys Ala Gly Gly Cys Ser Gly Pro Lys Asp Thr Asp Cys Phe Ala Cys Met Asn Phe Asn Asp Ser Gly Ala Cys Val Thr Gln Cys Pro Gln Thr Phe Val Tyr Asn Pro Thr Thr Phe Gln Leu Glu His Asn Phe Asn Ala Lys Tyr Thr Tyr Gly Ala Phe Cys Val Lys Lys Cys Pro His Asn Phe Val Val Asp Ser Ser Ser Cys Val Arg Ala Cys Pro Ser Ser Lys Met Glu Val Glu Glu Asn Gly Ile Lys Met Cys Lys Pro Cys Thr Asp Ile Cys Pro Lys Ala Cys Asp Gly Ile Gly Thr Gly Ser Leu Met Ser Ala Gln Thr Val Asp Ser Ser Asn Ile Asp Lys Phe Ile Asn Cys Thr Lys Ile 

Asn Gly Asn Leu Ile Phe Leu Val Thr Gly Ile His Gly Asp Pro Tyr

Asn Ala Ile Asp Ala Ile Asp Pro Glu Lys Leu Asn Val Phe Arg Thr Val Arg Glu Ile Thr Gly Phe Leu Asn Ile Gln Ser Trp Pro Pro Asn Met Thr Asp Phe Ser Val Phe Ser Asn Leu Val Thr Ile Gly Gly Arg Val Leu Tyr Ser Gly Leu Ser Leu Leu Ile Leu Lys Gln Gln Gly Ile Thr Ser Leu Gln Phe Gln Ser Leu Lys Glu Ile Ser Ala Gly Asn Ile Tyr Ile Thr Asp Asn Ser Asn Leu Cys Tyr Tyr His Thr Ile Asn Trp Thr Thr Leu Phe Ser Thr Ile Asn Gln Arg Ile Val Ile Arg Asp Asn Arg Arg Ala Glu Asn Cys Thr Ala Glu Gly Met Val Cys Asn His Leu Cys Ser Asn Asp Gly Cys Trp Gly Pro Gly Pro Asp Gln Cys Leu Ser Cys Arg Arg Phe Ser Arg Gly Lys Ile Cys Ile Glu Ser Cys Asn Leu Tyr Asp Gly Glu Phe Arg Glu Phe Glu Asn Gly Ser Ile Cys Val Glu Cys Asp Ser Gln Cys Glu Lys Met Glu Asp Gly Leu Leu Thr Cys His Gly Pro Gly Pro Asp Asn Cys Thr Lys Cys Ser His Phe Lys Asp Gly Pro Asn Cys Val Glu Lys Cys Pro Asp Gly Leu Gln Gly Ala Asn Ser 

Phe Ile Phe Lys Tyr Ala Asp Gln Asp Arg Glu Cys His Pro Cys His

Pro Asn Cys Thr Gln Gly Cys Asn Gly Pro Thr Ser His Asp Cys Ile 610 615 620

Tyr Tyr Pro Trp Thr Gly His Ser Thr Leu Pro Gln His Ala Arg Thr 625 630 635 640

Pro Leu Ile Ala Ala Gly Val Ile Gly Gly Leu Phe Ile Leu Val Ile 645 650 655

Met Ala Leu Thr Phe Ala Val Tyr Val Arg Arg Lys Ser Ile Lys Lys 660 665 670

Lys Arg Ala Leu Arg Arg Phe Leu Glu Thr Glu Leu Val Glu Pro Leu 675 680 685

Thr Pro Ser Gly Thr Ala Pro Asn Gln Ala Gln Leu Arg Ile Leu Lys 690 695 700

Glu Thr Glu Leu Lys Arg Val Lys Val Leu Gly Ser Gly Ala Phe Gly 705 710 715 720

Thr Val Tyr Lys Gly Ile Trp Val Pro Glu Gly Glu Thr Val Lys Ile 725 730 735

Pro Val Ala Ile Lys Ile Leu Asn Glu Thr Thr Gly Pro Lys Ala Asn 740 745 750

Val Glu Phe Met Asp Glu Ala Leu Ile Met Ala Ser Met Asp His Pro 755 760 765

His Leu Val Arg Leu Leu Gly Val Cys Leu Ser Pro Thr Ile Gln Leu 770 775 780

Val Thr Gln Leu Met Pro His Gly Cys Leu Leu Asp Tyr Val His Glu 785 790 795 800

His Lys Asp Asn Ile Gly Ser Gln Leu Leu Leu Asn Trp Cys Val Gln 805 810 815

Ile Ala Lys Gly Met Met Tyr Leu Glu Glu Arg Arg Leu Val His Arg 820 825 830

Asp Leu Ala Ala Arg Asn Val Leu Val Lys Ser Pro Asn His Val Lys 835 840 845

- Ile Thr Asp Phe Gly Leu Ala Arg Leu Leu Glu Gly Asp Glu Lys Glu 850 855 860
- Tyr Asn Ala Asp Gly Gly Lys Met Pro Ile Lys Trp Met Ala Leu Glu 865 870 875 880
- Cys Ile His Tyr Arg Lys Phe Thr His Gln Ser Asp Val Trp Ser Tyr 885 890 895
- Gly Val Thr Ile Trp Glu Leu Met Thr Phe Gly Gly Lys Pro Tyr Asp 900 905 910
- Gly Ile Pro Thr Arg Glu Ile Pro Asp Leu Leu Glu Lys Gly Glu Arg 915 920 925
- Leu Pro Gln Pro Pro Ile Cys Thr Ile Asp Val Tyr Met Val Met Val 930 935 940
- Lys Cys Trp Met Ile Asp Ala Asp Ser Arg Pro Lys Phe Lys Glu Leu 945 950 955 960
- Ala Ala Glu Phe Ser Arg Met Ala Arg Asp Pro Gln Arg Tyr Leu Val 965 970 975
- Ile Gln Gly Asp Asp Arg Met Lys Leu Pro Ser Pro Asn Asp Ser Lys 980 985 990
- Phe Phe Gln Asn Leu Leu Asp Glu Glu Asp Leu Glu Asp Met Met Asp 995 1000 1005
- Ala Glu Glu Tyr Leu Val Pro Gln Ala Phe Asn Ile Pro Pro Pro 1010 1015 1020
- Ile Tyr Thr Ser Arg Thr Arg Ile Asp Ser Asn Arg Ser Glu Ile 1025 1030 1035
- Gly His Ser Pro Pro Pro Ala Tyr Thr Pro Met Ser Gly Asn Gln 1040 1045 1050
- Phe Val Tyr Gln Asp Gly Gly Phe Ala Thr Gln Gly Met Pro 1055 1060 1065
- Met Pro Tyr Arg Ala Thr Thr Ser Thr Ile Pro Glu Ala Pro Val 1070 1075 1080

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Ala Gln Gly Ala Thr Ala Glu Met Phe Asp Asp Ser Cys Cys Asn Gly Thr Leu Arg Lys Pro Val Ala Pro His Val Gln Glu Asp Ser Ser Thr Gln Arg Tyr Ser Ala Asp Pro Thr Val Phe Ala Pro Glu Arg Asn Pro Arg Gly Glu Leu Asp Glu Gly Tyr Met Thr Pro Met His Asp Lys Pro Lys Gln Glu Tyr Leu Asn Pro Val Glu Glu Asn Pro Phe Val Ser Arg Arg Lys Asn Gly Asp Leu Gln Ala Leu Asp Asn Pro Glu Tyr His Ser Ala Ser Ser Gly Pro Pro Lys Ala Glu Asp Glu Tyr Val Asn Glu Pro Leu Tyr Leu Asn Thr Phe Ala Asn Ala Leu Gly Ser Ala Glu Tyr Met Lys Asn Ser Val Leu Ser Val Pro Glu Lys Ala Lys Lys Ala Phe Asp Asn Pro Asp Tyr Trp Asn His Ser Leu Pro Pro Arg Ser Thr Leu Gln His Pro Asp Tyr Leu Gln Glu Tyr Ser Thr Lys Tyr Phe Tyr Lys Gln Asn Gly Arg Ile Arg Pro Ile Val Ala Glu Asn Pro Glu Tyr Leu Ser Glu Phe Ser Leu Lys Pro Gly Thr Met Leu Pro Pro Pro Pro Tyr Arg His Arg Asn Thr Val Val